

RECEIVED Page 1 of 7
#8

NOV 13 2001

TECH CENTER 1600/2900

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/770,643A

DATE: 09/10/2001

TIME: 08:33:19

Input Set : A:\LEX-0122-USA SEQLIST.txt

Output Set: N:\CRF3\09102001\I770643A.raw

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NOV 13 2001

TECH CENTER 1600/2900

4 <110> APPLICANT: Turner, C. Alexander Jr.
5 Hilbun, Erin
6 Donoho, Gregory
7 Scoville, John
8 Wattler, Frank
9 Friedrich, Glenn
10 Abuin, Alejandro
11 Zambrowicz, Brian
12 Sands, Arthur T.

14 <120> TITLE OF INVENTION: Novel Human Neurexin-like Proteins and Polynucleotides
Encoding the

15 Same
17 <130> FILE REFERENCE: LEX-0166-PRV
C--> 19 <140> CURRENT APPLICATION NUMBER: US/09/770,643A
C--> 19 <141> CURRENT FILING DATE: 2001-01-26
19 <160> NUMBER OF SEQ ID NOS: 31
21 <170> SOFTWARE: FastSEQ for Windows Version 4.0
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 3924
25 <212> TYPE: DNA
26 <213> ORGANISM: homo sapiens
28 <400> SEQUENCE: 1

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31 atggctttt ccagttcctc agacactcact ggcactcaca gcccagctca actcaactgg 180
32 agagttggaa ctggcggtt gtcggccatg gattccaatg ctcaacagtg gctccagatg 240
33 gacctgggaa acagagtaga gattacagca gtggccacgc agggaaagata cggaaagctct 300
34 gactgggtga cgagttacag cctgatgttc agtgcacacag gacgcaactg gaaacagttac 360
35 aaacaagaag acagcatctg gacctttgca gaaacatga atgctgacag cgtggtgac 420
36 cacaagctat tgcactcagt gagagcccgaa tttttcgct ttgtccccct ggaatggaaat 480
37 cccagtggaa agattggcat gagagtcgag gtctacggat gttcctataa atcagacgtt 540
38 gctgactttg atggccgaag ctcacttctg tacaggttca atcagaagtt gatgagttact 600
39 ctcaaagatg tgcactccct gaagttcaag agcatgcaag gagatgggt cctgttccat 660
40 ggagaaggta agcgtggaga ccacatcacc ttgaaactcc agaaggggag gctggcccta 720
41 cacctaatt tgggtgacag caaagcgcgg ctcagcagca gcttggccctc tgccaccctg 780
42 ggcagcctcc tggatgacca gcactggcac tyggcctca ttgagcgggt gggcaagcag 840
43 gtgaacttca cggtgacaa gcacacacag cacttccgca ccaaggcgaa gacggatgcc 900
44 ttagacattt actatgagct tagtttggaa ggaattccag taccaggaaa acctggacc 960
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46 attracctgg ctaagagacg aaagcatcag atctatactg tggcaatgt cacttttcc 1080
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48 ctgcccggca ccccccaat tggatggctc tcagttagtt tccagttcg aacatggaaac 1200
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53 gtgcagattt attctggaaa tagtactat tttggagggtt gccccgacaa tctcaccgat 1500
54 tcccaatgtt taaatcccat taaggcttc caaggctgca tgaggctcat ctttattgt 1560

ENTERED

A.S

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/770,643A

DATE: 09/10/2001
TIME: 08:33:49

Input Set : A:\LEX-0122-USA SEQLIST.txt
Output Set: N:\CRF3\09102001\I770643A.raw

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57	agctgctccc	agtccctggac	taccttctat	tgtaactgca	gtgacacaag	ttacactggt	1740
58	gccacctgcc	acaactccat	ctacgagcaa	tcctgcgagg	tgtacaggca	ccagggaaat	1800
59	acagccggct	tcttctacat	cgactcagat	ggcagcggcc	cactgggacc	tctccaggtg	1860
60	tactgcaata	tcaactgagga	caagatctgg	acatcagtgc	agcacaacaa	tacagagctg	1920
61	acccgagtgc	ggggcgctaa	ccctgagaag	ccctatgcca	tggcatttgg	ctacggggc	1980
62	agcatggaac	agctggaggc	cgtgatcgac	ggctctgagc	actgtgagca	ggaggtggcc	2040
63	taccactgca	ggaggtcccc	cctgctcaac	acgccggatg	gaacaccatt	tacctgggtgg	2100
64	attggcgggt	ccaatgaaag	gcacccttac	tggggaggtt	cccctcctgg	ggtccagcag	2160
65	tgtgagtgt	gcctagacga	gagctgcctg	gacattcagc	actttgcaa	ttgcgacgct	2220
66	gacaaggatg	aatggacaaa	tgatactggc	tttctttcct	tcaaagacca	cttgcctgtc	2280
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69	tcttacctcc	actttcctac	cttccatgct	gaattcagt	ccgatatttc	cttcttttt	2460
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97	<212>	TYPE:	PRT				
98	<213>	ORGANISM:	homo sapiens				
99	<220>	FEATURE:					
100	<221>	NAME/KEY:	VARIANT				
101	<222>	LOCATION:	(1)...(1307)				
102	<223>	OTHER INFORMATION:	Xaa = Any Amino Acid				
103	<400>	SEQUENCE:	2				

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107 Met Asp Ser Leu Pro Arg Leu Thr Ser Val Leu Thr Leu Leu Phe Ser
108 1 5 10 15
109 Gly Leu Trp His Leu Gly Leu Thr Ala Thr Asn Tyr Asn Cys Asp Asp
110 20 25 30
111 Pro Leu Ala Ser Leu Leu Ser Pro Met Ala Phe Ser Ser Ser Asp
112 35 40 45
113 Leu Thr Gly Thr His Ser Pro Ala Gln Leu Asn Trp Arg Val Gly Thr
114 50 55 60
115 Gly Gly Trp Ser Pro Ala Asp Ser Asn Ala Gln Gln Trp Leu Gln Met
116 65 70 75 80
117 Asp Leu Gly Asn Arg Val Glu Ile Thr Ala Val Ala Thr Gln Gly Arg
118 85 90 95
119 Tyr Gly Ser Ser Asp Trp Val Thr Ser Tyr Ser Leu Met Phe Ser Asp
120 100 105 110
121 Thr Gly Arg Asn Trp Lys Gln Tyr Lys Gln Glu Asp Ser Ile Trp Thr
122 115 120 125
123 Phe Ala Gly Asn Met Asn Ala Asp Ser Val Val His His Lys Leu Leu
124 130 135 140
125 His Ser Val Arg Ala Arg Phe Val Arg Phe Val Pro Leu Glu Trp Asn
126 145 150 155 160
127 Pro Ser Gly Lys Ile Gly Met Arg Val Glu Val Tyr Gly Cys Ser Tyr
128 165 170 175
129 Lys Ser Asp Val Ala Asp Phe Asp Gly Arg Ser Ser Leu Leu Tyr Arg
130 180 185 190
131 Phe Asn Gln Lys Leu Met Ser Thr Leu Lys Asp Val Ile Ser Leu Lys
132 195 200 205
133 Phe Lys Ser Met Gln Gly Asp Gly Val Leu Phe His Gly Glu Gly Gln
134 210 215 220
135 Arg Gly Asp His Ile Thr Leu Glu Leu Gln Lys Gly Arg Leu Ala Leu
136 225 230 235 240
137 His Leu Asn Leu Gly Asp Ser Lys Ala Arg Leu Ser Ser Ser Leu Pro
138 245 250 255
W--> 139 Ser Ala Thr Leu Gly Ser Leu Leu Asp Asp Gln His Trp His Xaa Val
140 260 265 270
141 Leu Ile Glu Arg Val Gly Lys Gln Val Asn Phe Thr Val Asp Lys His
142 275 280 285
143 Thr Gln His Phe Arg Thr Lys Gly Glu Thr Asp Ala Leu Asp Ile Asp
144 290 295 300
145 Tyr Glu Leu Ser Phe Gly Gly Ile Pro Val Pro Gly Lys Pro Gly Thr
146 305 310 315 320
147 Phe Leu Lys Lys Asn Phe His Gly Cys Ile Glu Asn Leu Tyr Tyr Asn
148 325 330 335
W--> 149 Gly Val Asn Ile Ile Xaa Leu Ala Lys Arg Arg Lys His Gln Ile Tyr
150 340 345 350
151 Thr Val Gly Asn Val Thr Phe Ser Cys Ser Glu Pro Gln Ile Val Pro
152 355 360 365
153 Ile Thr Phe Val Asn Ser Ser Gly Ser Tyr Leu Leu Leu Pro Gly Thr
154 370 375 380
155 Pro Gln Ile Asp Gly Leu Ser Val Ser Phe Gln Phe Arg Thr Trp Asn

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156	385	390	395	400
157	Lys Asp Gly Leu Leu Leu Ser Thr Glu Leu Ser Glu Gly Ser Gly Thr			
158		405	410	415
159	Leu Leu Leu Ser Leu Glu Gly Gly Ile Leu Arg Leu Val Ile Gln Lys			
160		420	425	430
161	Met Thr Glu Arg Val Ala Glu Ile Leu Thr Gly Ser Asn Leu Asn Asp			
162		435	440	445
163	Gly Leu Trp His Ser Val Ser Ile Asn Ala Arg Arg Asn Arg Ile Thr			
164		450	455	460
165	Leu Thr Leu Asp Asp Glu Ala Ala Pro Pro Ala Pro Asp Ser Thr Trp			
166		465	470	475
167	Val Gln Ile Tyr Ser Gly Asn Ser Tyr Tyr Phe Gly Gly Cys Pro Asp			
168		485	490	495
169	Asn Leu Thr Asp Ser Gln Cys Leu Asn Pro Ile Lys Ala Phe Gln Gly			
170		500	505	510
171	Cys Met Arg Leu Ile Phe Ile Asp Asn Gln Pro Lys Asp Leu Ile Ser			
172		515	520	525
173	Val Gln Gln Gly Ser Leu Gly Asn Phe Ser Asp Leu His Ile Asp Leu			
174		530	535	540
175	Cys Ser Ile Lys Asp Arg Cys Leu Pro Asn Tyr Cys Glu His Gly Gly			
176		545	550	555
177	560			
178	Ser Cys Ser Gln Ser Trp Thr Thr Phe Tyr Cys Asn Cys Ser Asp Thr			
179		565	570	575
180	Ser Tyr Thr Gly Ala Thr Cys His Asn Ser Ile Tyr Glu Gln Ser Cys			
181		580	585	590
182	Glu Val Tyr Arg His Gln Gly Asn Thr Ala Gly Phe Phe Tyr Ile Asp			
183		595	600	605
184	Ser Asp Gly Ser Gly Pro Leu Gly Pro Leu Gln Val Tyr Cys Asn Ile			
185		610	615	620
186	Thr Glu Asp Lys Ile Trp Thr Ser Val Gln His Asn Asn Thr Glu Leu			
187		625	630	635
188	640			
189	Thr Arg Val Arg Gly Ala Asn Pro Glu Lys Pro Tyr Ala Met Ala Leu			
190		645	650	655
191	Asp Tyr Gly Gly Ser Met Glu Gln Leu Glu Ala Val Ile Asp Gly Ser			
192		660	665	670
193	Glu His Cys Glu Gln Glu Val Ala Tyr His Cys Arg Arg Ser Arg Leu			
194		675	680	685
195	690			
196	Leu Asn Thr Pro Asp Gly Thr Pro Phe Thr Trp Trp Ile Gly Arg Ser			
197		695	700	
198	710			
199	Asn Glu Arg His Pro Tyr Trp Gly Gly Ser Pro Pro Gly Val Gln Gln			
200		715		
201	720			
202	Cys Glu Cys Gly Leu Asp Glu Ser Cys Leu Asp Ile Gln His Phe Cys			
203		725	730	735
204	Asn Cys Asp Ala Asp Lys Asp Glu Trp Thr Asn Asp Thr Gly Phe Leu			
		740	745	750
205	Ser Phe Lys Asp His Leu Pro Val Thr Gln Ile Val Ile Thr Asp Thr			
206		755	760	765
207	Asp Arg Ser Asn Ser Glu Ala Ala Trp Arg Ile Gly Pro Leu Arg Cys			
208		770	775	780

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205 Tyr Gly Asp Arg Arg Phe Trp Asn Ala Val Ser Phe Tyr Thr Glu Ala
206 785 790 795 800
207 Ser Tyr Leu His Phe Pro Thr Phe His Ala Glu Phe Ser Ala Asp Ile
208 805 810 815
209 Ser Phe Phe Phe Lys Thr Thr Ala Leu Ser Gly Val Phe Leu Glu Asn
210 820 825 830
211 Leu Gly Ile Lys Asp Phe Ile Arg Leu Glu Ile Ser Ser Pro Ser Glu
212 835 840 845
213 Ile Thr Phe Ala Ile Asp Val Gly Asn Gly Pro Val Glu Leu Val Val
214 850 855 860
215 Gln Ser Pro Ser Leu Leu Asn Asp Asn Gln Trp His Tyr Val Arg Ala
216 865 870 875 880
217 Glu Arg Asn Leu Lys Glu Thr Ser Leu Gln Val Asp Asn Leu Pro Arg
218 885 890 895
219 Ser Thr Arg Glu Thr Ser Glu Glu Gly His Phe Arg Leu Gln Leu Asn
220 900 905 910
221 Ser Gln Leu Phe Val Gly Gly Thr Ser Ser Arg Gln Lys Gly Phe Leu
222 915 920 925
223 Gly Cys Ile Arg Ser Leu His Leu Asn Gly Gln Lys Met Asp Leu Glu
224 930 935 940
225 Glu Arg Ala Lys Val Thr Ser Gly Val Arg Pro Gly Cys Pro Gly His
226 945 950 955 960
227 Cys Ser Ser Tyr Gly Ser Ile Cys His Asn Gly Gly Lys Cys Val Glu
228 965 970 975
229 Lys His Asn Gly Tyr Leu Cys Asp Cys Thr Asn Ser Pro Tyr Glu Gly
230 980 985 990
231 Pro Phe Cys Lys Lys Glu Val Ser Ala Val Phe Glu Ala Gly Thr Ser
232 995 1000 1005
233 Val Thr Tyr Met Phe Gln Glu Pro Tyr Pro Val Thr Lys Asn Ile Ser
234 1010 1015 1020
235 Leu Ser Ser Ser Ala Ile Tyr Thr Asp Ser Ala Pro Ser Lys Glu Asn
236 1025 1030 1035 1040
237 Ile Ala Leu Ser Phe Val Thr Thr Gln Ala Pro Ser Leu Leu Phe
238 1045 1050 1055
239 Ile Asn Ser Ser Ser Gln Asp Phe Val Val Leu Leu Cys Lys Asn
240 1060 1065 1070
241 Gly Ser Leu Gln Val Arg Tyr His Leu Asn Lys Glu Glu Thr His Val
242 1075 1080 1085
243 Phe Thr Ile Asp Ala Asp Asn Phe Ala Asn Arg Arg Met His His Leu
244 1090 1095 1100
245 Lys Ile Asn Arg Glu Gly Arg Glu Leu Thr Ile Gln Met Asp Gln Gln
246 1105 1110 1115 1120
247 Leu Arg Leu Ser Tyr Asn Phe Ser Pro Glu Val Glu Phe Arg Val Ile
248 1125 1130 1135
249 Arg Ser Leu Thr Leu Gly Lys Val Thr Glu Asn Leu Gly Leu Asp Ser
250 1140 1145 1150
251 Glu Val Ala Lys Ala Asn Ala Met Gly Phe Ala Gly Cys Met Ser Ser
252 1155 1160 1165
253 Val Gln Tyr Asn His Ile Ala Pro Leu Lys Ala Ala Leu Arg His Ala

Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/770,643A

DATE: 09/10/2001

TIME: 08:33:50

Input Set : A:\LEX-0122-USA SEQLIST.txt
Output Set: N:\CRF3\09102001\I770643A.raw

L:19 M:270 C: Current Application Number differs, Replaced Current Application No
L:19 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:139 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:149 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:385 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:657 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:742 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:752 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:862 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:987 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:997 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:1135 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:1284 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:1294 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:1449 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22